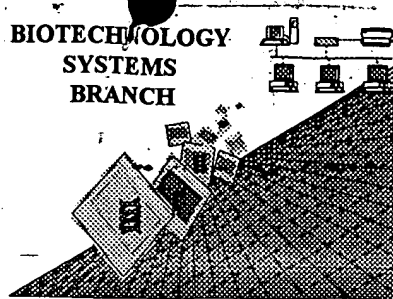


RAW SEQUENCE LISTING **ERROR REPORT**

BIOTECHNOLOGY
SYSTEMS
BRANCH



Handwritten: 0400
1-23-01

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/745,167

Source: OIPE

Date Processed by STIC: 1/10/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/745,167

DATE: 01/10/2001
TIME: 12:24:42

Input Set : A:\RTS-0140 Sequence Listing.txt
Output Set: N:\CRF3\01102001\I745167.raw

**Does Not Comply
Corrected Diskette Needed**

OK
6 <110> APPLICANT: Brett P. Monia
7 Jacqueline Wyatt
9 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF HISTONE DEACETYLASE 1 EXPRESSION
11 <130> FILE REFERENCE: RTS-0140
13 <140> CURRENT APPLICATION NUMBER: US/09/745,167
13 <141> CURRENT FILING DATE: 2000-12-19
13 <160> NUMBER OF SEQ ID NOS: 87

ERRORED SEQUENCES

264 <210> SEQ ID NO: 9
265 <211> LENGTH: 22
266 <212> TYPE: DNA
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
271 <223> OTHER INFORMATION: PCR Probe
273 <400> SEQUENCE: 9
E--> 274 cgccgtgtca ccagggctgc tx

22

*do not use "x" in the sequence itself, use "n" and
explain in <220>-<223> section*

VERIFICATION SUMMARY DATE: 01/10/2001
PATENT APPLICATION: US/09/745,167 TIME: 12:24:43

Input Set : A:\RTS-0140 Sequence Listing.txt
Output Set: N:\CRF3\01102001\I745167.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:274 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:1292 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:3
L:1295 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:87
L:1295 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:3
L:1299 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:87
L:1299 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:4
L:1300 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:87
L:1300 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:4